



```

chain nodes :
  23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 47 48
ring nodes :
  1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22
ring/chain nodes :
  44
chain bonds :
  1-10 2-31 3-32 4-24 5-33 6-34 9-37 11-23 13-35 14-36 15-25 16-22 17-26 17-39
  18-27 18-40 19-28 19-41 20-29 20-42 22-38 24-47 25-48 29-30 43-44
ring bonds :
  1-2 1-6 2-3 3-4 4-5 5-6 7-8 7-12 7-16 8-9 9-10 10-11 11-12 12-13 13-14 14-15
  15-16 17-18 17-22 18-19 19-20 20-21 21-22
exact/norm bonds :
  4-24 7-8 8-9 9-10 10-11 11-12 11-23 15-25 17-18 17-22 17-26 18-19 18-27 19-20
  19-28 20-21 21-22 24-47 25-48 29-30 43-44
exact bonds :
  1-10 2-31 3-32 5-33 6-34 9-37 13-35 14-36 16-22 17-39 18-40 19-41 20-29 20-42
  22-38
normalized bonds :
  1-2 1-6 2-3 3-4 4-5 5-6 7-12 7-16 12-13 13-14 14-15 15-16

```

G1:H, [*1]

```

Match level :
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom
12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom
22:Atom 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS
31:CLASS 32:CLASS 33:CLASS 34:CLASS 35:CLASS 36:CLASS 37:CLASS 38:CLASS 39:CLASS
40:CLASS 41:CLASS 42:CLASS 43:CLASS 44:CLASS 47:CLASS 48:CLASS

```